## REPLACEMENT SHEET

## FIGURE 6: 5/44 HEAVY CHAIN SEQUENCE GRAFT DESIGN

|             | 10                            | 20         | 30       |        | 40                  | 50              |    |
|-------------|-------------------------------|------------|----------|--------|---------------------|-----------------|----|
| $V_{H}$     | EVQLQQSGTVLARI                | PGASVKMSCK | ASGYRFT  | NYWIH  | WVKQRPGQG:          | LEWIG GI        | ΝP |
|             |                               | 1          |          |        |                     |                 |    |
| DP7         | QVQLVQSGAEVKKI                | PGASVKVSCK | ASGYTFT  |        | WVRQAPGQG           | LEWMG           |    |
|             |                               |            |          |        |                     |                 |    |
| gH1         | EVQLVQSGAEVKK                 | PGASVKVSCK | ASGYRFT  | NYWIH  | WVRQAPGQG           | LEWIG <b>GI</b> | ΝP |
| gH4,5,6,7   | EVQLVQSGAEVKK                 | PGASVKVSCK | ASGYRFT  | NYWIH  | WVRQAPGQG           | LEWIG <i>GI</i> | ΝP |
|             |                               |            |          |        |                     |                 |    |
|             | 60                            | 70         | 80       |        | 90                  | 100             |    |
| $V_H$       | GNNYTTYKRNLKG                 | KATLTAVTS  | ASTAYMDL | SSLTSE | EDSAVYYCTR          | EGYGNYG         |    |
|             |                               |            |          |        |                     |                 |    |
| DP7         | KFQG                          | RVTMTRDTS  | TSTVYMEL | SSLRSE | EDTAVYYCAR          |                 |    |
|             |                               |            |          |        |                     |                 |    |
| gH1         | GNQYTTYKRNLKG                 |            |          |        |                     |                 |    |
| gH4         | GNNYATYRRNLKG                 |            |          |        |                     |                 |    |
| gH5         | GNNYATYRRNLKG                 | _          |          |        | _                   |                 |    |
| gH6         | GNNYATYRRKF QG                |            |          |        |                     |                 |    |
| gH7         | <b>GNNYATYRR</b> KF <b>QG</b> | RVTMTADTS  | TSTVYMEL | SSLRSE | EDTAVYYC <u>T</u> R | EGYGNYG         |    |
|             |                               |            |          |        |                     |                 |    |
|             | 110                           |            |          |        |                     |                 |    |
| JH4         | WGQGTLV'                      | rvss       |          |        |                     |                 |    |
|             |                               |            |          |        |                     |                 |    |
| $V_{\rm H}$ | AWFAY WGQGTLV                 | TVSS       |          |        |                     |                 |    |
|             |                               |            |          |        |                     |                 |    |
| gHl         | AWFAY WGQGTLVTVSS             |            |          |        |                     |                 |    |
| gH4,5,6,7   | AWFAY WGQGTLV                 | rvss       |          |        |                     |                 |    |